SEQUENCE LISTING

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       Rebecca E. Cahoon
       William D. Hitz
       Anthony J. Kinney
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310

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Gly Leu His Phe Gln Ile Glu His His Leu Phe Pro Arg Met Pro Lys 380 375

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285

280

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Phe Trp Ala Trp Tyr Pro Leu Leu Val Ala Ser Leu Pro Asn Trp Trp 305 310 315 320

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His Val Gln Phe Cys Leu Asn His Phe Ser Ser Asp Val Tyr Val Gly 340 345 350

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Phe Gln Ile Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu 385 390 395 400

Arg Lys Val Ala Pro Ala Val Arg Asp Leu Cys Lys Lys His Gly Leu 405 410 415

Thr Tyr Ser Ala Ala Thr Phe Trp Gly Ala Asn Val Leu Thr Trp Lys 420 425 430

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Ser Thr Phe Val His Val Leu Ser Ala Ala Leu Ile Gly Phe Leu Trp 70 Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His Tyr Asn Val Met 90 Leu Ser Arg Arg Leu Asn Arg Ala Ile Gln Ile Leu Ser Gly Asn Ile 105 Leu Ala Gly Ile Ser Ile Gly Trp Trp Lys Trp Asn His Asn Ala His 120 His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Met 135 Pro Val Phe Ala Val Ser Ser Arg Phe Phe Asn Ser Ile Thr Ser His 155 Xaa Tyr Gly Arg Lys Xaa Glu Phe Asp Xaa Ile Ala Xaa Phe Leu Ile 170 Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Phe Ser Arg Xaa Lys Val 200 Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp 215 Phe Leu Phe Leu Leu Ala Leu Leu Phe Val Pro Ile Gln His Ile Gln 235 230 Phe Trp Leu Asn His Leu Ala Glu Asn Leu Tyr Xaa Gly 245

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Val Tyr Asn Val Ser Asp Trp Val Lys Glu His Pro Gly Gly Asp Val 35 40 45

Pro Ile Ser Asn Leu Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala 50 55 60

Tyr His Pro Gly Thr Ala Trp Ser His Leu Glu Lys Phe Phe Thr Gly 65 70 75 80

Tyr His Leu Ser Asp Phe Lys Val Ser Glu Val Ser Lys Asp Tyr Arg 85 90 95

Lys Leu Ala Ser Glu Phe Ser Lys Leu Gly Leu Phe Asp Thr Lys Gly 100 105 110

His Val Thr Ser Cys Thr Leu Ala Ser Val Ala Val Met Phe Leu Ile 115 120 125

Val Leu Tyr Gly Val Leu Arg Cys Thr Ser Val Trp Ala His Leu Gly 130 135 140

Ser Gly Met Leu Leu Gly Leu Leu Trp Met Gln Ser Ala Tyr Val Gly 145 150 155 160

His Asp Ser Gly His Tyr Val Val Met Thr Thr Asn Gly Phe Asn Lys
165 170 175

Val Ala Gln Ile Leu Ser Gly Asn Cys Leu Thr Gly Ile Ser Ile Ala 180 185 190

Trp Trp Lys Trp Thr His Asn Ala His His Ile Ala Cys Asn Ser Leu 195 200 205

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Tyr	Pro	Val	Met 260	Cys	Val	Ala	Arg	Val 265	Asn	Leu	Tyr	Leu	Gln 270	Thr	Ile	
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Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly

Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala

Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe

Phe Val Gly Arg Leu Ser Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp 105

Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg 115 120

Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe 130 135

Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His

145					150					155					160
Met	Phe	Ala	Gly	Gly 165	Leu	Ile	Gly	Phe	Ile 170	Trp	Ile	Gln	Ser	Gly 175	Trp
Île	Gly	His	Asp 180	Ser	Gly	His	His	Gln 185	Ile	Thr	Arg	His	Pro 190	Ala	Leu
Asn	Arg	Leu 195	Leu	Gln	Val	Val	Ser 200	Gly	Asn	Cys	Leu	Thr 205	Gly	Leu	Gly
Ile	Ala 210	Trp	Trp	Lys	Phe	Asn 215	His	Asn	Thr	His	His 220	Ile	Ser	Cys	Asn
Ser 225	Leu	Asp	His	Asp	Pro 230	Asp	Leu	Gln	His	Leu 235	Pro	Leu	Phe	Ala	Val 240
Ser	Thr	Lys	Leu	Phe 245	Asn	Asn	Leu	Trp	Ser 250	Val	Суѕ	Tyr	Glu	Arg 255	Thr
Leu	Ala	Phe	Asp 260	Ala	Ile	Ser	Lys	Phe 265	Phe	Val	Ser	Tyr	Gln 270	His	Trp
Thr	Phe	Tyr 275	Pro	Val	Met	Gly	Phe 280	Ala	Arg	Ile	Asn	Leu 285	Leu	Val	Gln
Ser	Ile 290	Val	Phe	Leu	Ile	Thr 295	Gln	Lys	Lys	Val	Arg 300	Gln	Arg	Trp	Leu
Glu 305	Ile	Ala	Gly	Val	Ala 310	Ala	Phe	Trp	Val	Trp 315	Tyr	Pro	Leu	Leu	Val 320
Ser	Cys	Leu	Pro	Asn 325	Trp	Trp	Glu	Arg	Val 330	Ala	Phe	Val	Leu	Ala 335	Ser
Phe	Val	Ile	Thr 340	Gly	Ile	Gln	His	Val 345	Gln	Phe	Cys	Leu	Asn 350	His	Phe
Ser	Ser	Ala 355	Val	Tyr	Val	Gly	Pro 360	Pro	Lys	Gly	Asn	Asp 365	Trp	Phe	Glu
Arg	Gln 370	Thr	Ala	Gly	Thr	Leu 375	Asp	Ile	Lys	Cys	Ser 380	Pro	Trp	Met	Asp
Trp 385	Phe	His	Gly	Gly	Leu 390	Gln	Phe	Gln	Val	Glu 395	His	His	Leu	Phe	Pro 400
Arg	Leu	Pro	Arg	Cys 405	His	Tyr	Arg	Met	Val 410	Ala	Pro	Ile	Val	Arg 415	Asp
Leu	Cys	Lys	Lys 420	His	Gly	Leu	Ser	Tyr 425	Gly	Ala	Ala	Thr	Phe 430	Trp	Glu
Ala	Asn	Val 435	Met	Thr	Trp	Lys	Thr 440	Leu	Arg	Ala	Ala	Ala 445	Leu	Gln	Ala
Arg	Glu 450	Ala	Thr	Thr	Gly	Ala 455	Ala	Pro	Lys	Asn	Leu 460	Val	Trp	Glu	Ala
Leu 465	Asn	Thr	His	Gly											

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His Asp Lys Pro Gly Asp Leu Trp Ile Ser Ile Gln Gly Lys Ala Tyr 20 25 30

Asp Val Ser Asp Trp Val Lys Asp His Pro Gly Gly Ser Phe Pro Leu 35 40 45

Lys Ser Leu Ala Gly Gln Glu Val Thr Asp Ala Phe Val Ala Phe His
50 55 60

Pro Ala Ser Thr Trp Lys Asn Leu Asp Lys Phe Phe Thr Gly Tyr Tyr 65 70 75 80

Leu Lys Asp Tyr Ser Val Ser Glu Val Ser Lys Asp Tyr Arg Lys Leu 85 90 95

Val Phe Glu Phe Ser Lys Met Gly Leu Tyr Asp Lys Lys Gly His Ile 100 105 110

Met Phe Ala Thr Leu Cys Phe Ile Ala Met Leu Phe Ala Met Ser Val 115 120 125

Tyr Gly Val Leu Phe Cys Glu Gly Val Leu Val His Leu Phe Ser Gly 130 135 140

Cys Leu Met Gly Phe Leu Trp Ile Gln Ser Gly Trp Ile Gly His Asp 145 150 155 160

Ala Gly His Tyr Met Val Val Ser Asp Ser Arg Leu Asn Lys Phe Met 165 170 175

Gly Ile Phe Ala Ala Asn Cys Leu Ser Gly Ile Ser Ile Gly Trp Trp 180 185 190

Lys Trp Asn His Asn Ala His His Ile Ala Cys Asn Ser Leu Glu Tyr 195 200 205

Asp Pro Asp Leu Gln Tyr Ile Pro Phe Leu Val Val Ser Ser Lys Phe 210 215 220

Phe Gly Ser Leu Thr Ser His Phe Tyr Glu Lys Arg Leu Thr Phe Asp 225 230 235 240

Ser Leu Ser Arg Phe Phe Val Ser Tyr Gln His Trp Thr Phe Tyr Pro 245 250 255

Ile Met Cys Ala Ala Arg Leu Asn Met Tyr Val Gln Ser Leu Ile Met 260 265 270

Leu Leu Thr Lys Arg Asn Val Ser Tyr Arg Ala His Glu Leu Leu Gly 275 280 285

Cys Leu Val Phe Ser Ile Trp Tyr Pro Leu Leu Val Ser Cys Leu Pro 295 Asn Trp Gly Glu Arg Ile Met Phe Val Ile Ala Ser Leu Ser Val Thr Gly Met Gln Gln Val Gln Phe Ser Leu Asn His Phe Ser Ser Val Tyr Val Gly Lys Pro Lys Gly Asn Asn Trp Phe Glu Lys Gln Thr Asp 345 Gly Thr Leu Asp Ile Ser Cys Pro Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Ile Glu His His Leu Phe Pro Lys Met Pro Arg 375 Cys Asn Leu Arg Lys Ile Ser Pro Tyr Val Ile Glu Leu Cys Lys His Asn Leu Pro Tyr Asn Tyr Ala Ser Phe Ser Lys Ala Asn Glu Met 410 Thr Leu Arg Thr Leu Arg Asn Thr Ala Leu Gln Ala Arg Asp Ile Thr Lys Pro Leu Pro Lys Asn Leu Val Trp Glu Ala Leu His Thr His Gly 440

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Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly

Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala

Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe

Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp 105

Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg 120 115

Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe



						_									
Cys 145	Ile	Ala	Leu	Tyr	Cys 150	Val	Leu	Ala	Суз	Ser 155	Ser	Thr	Gly	Ala	His 160
Met	Phe	Ala	Gly	Gly 165	Leu	Ile	Gly	Phe	Ile 170	Trp	Ile	Gln	Ser	Gly 175	Trp
Ile	Gly	His	Asp 180	Ser	Gly	His	His	Gln 185	Ile	Thr	Arg	His	Pro 190	Ala	Leu
Asn	Arg	Leu 195	Leu	Gln	Val	Val	Ser 200	Gly	Asn	Cys	Leu	Thr 205	Gly	Leu	Gly
Ile	Ala 210	Trp	Trp	Lys	Phe	Asn 215	His	Asn	Thr	His	His 220	Ile	Ser	Cys	Asn
Ser 225	Leu	Asp	His	Asp	Pro 230	Asp	Leu	Gln	His	Leu 235	Pro	Leu	Phe	Ala	Val 240
Ser	Thr	Lys	Leu	Phe 245	Asn	Asn	Leu	Trp	Ser 250	Val	Cys	Tyr	Glu	Arg 255	Thr
Leu	Ala	Phe	Asp 260	Ala	Ile	Ser	Lys	Phe 265	Phe	Val	Ser	Tyr	Gln 270	His	Trp
Thr	Phe	Tyr 275	Pro	Val	Met	Gly	Phe 280	Ala	Arg	Ile	Asn	Leu 285	Leu	Val	Gln
Ser	Ile 290	Val	Phe	Leu	Ile	Thr 295	Gln	Lys	Lys	Val	Arg 300	Gln	Arg	Trp	Leu
Glu 305	Ile	Ala	Gly	Val	Ala 310	Ala	Phe	Trp	Val	Trp 315	Tyr	Pro	Leu	Leu	Val 320
Ser	Cys	Leu	Pro	Asn 325	Trp	Trp	Glu	Arg	Val 330	Ala	Phe	Val	Leu	Ala 335	Ser
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Arg	Gln 370	Thr	Ala	Gly	Thr	Leu 375	Asp	Ile	Lys	Суѕ	Ser 380	Pro	Trp	Met	Asp
Trp 385	Phe	His	Gly	Gly	Leu 390	Gln	Phe	Gln	Val	Glu 395	His	His	Leu	Phe	Pro 400
Arg	Leu	Pro	Arg	Cys 405	His	Tyr	Arg	Met	Val 410	Ala	Pro	Ile	Val	Arg 415	Asp
Leu	Cys	Lys	Lys 420	His	Gly	Leu	Ser	Tyr 425	Gly	Ala	Ala	Thr	Phe 430	Trp	Glu
Ala	Asn	Val 435	Met	Thr	Trp	Lys	Thr 440	Leu	Arg	Ala	Ala	Ala 445	Leu	Gln	Ala
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Lys Tyr Ile Thr Ser Lys Glu Leu Lys Lys His Asn Asn Pro Asn Asp 20 25 30

Leu Trp Ile Ser Ile Leu Gly Lys Val Tyr Asn Val Thr Glu Trp Ala 35 40 45

Lys Glu His Pro Gly Gly Asp Ala Pro Leu Ile Asn Leu Ala Gly Gln 50 55 60

Asp Val Thr Asp Ala Phe Ile Ala Phe His Pro Gly Thr Ala Trp Lys
65 70 75 80

His Leu Asp Lys Leu Phe Thr Gly Tyr His Leu Lys Asp Tyr Gln Val 85 90 95

Ser Asp Ile Ser Arg Asp Tyr Arg Lys Leu Ala Ser Glu Phe Ala Lys 100 105 110

Ala Gly Met Phe Glu Lys Lys Gly His Gly Val Ile Tyr Ser Leu Cys 115 120 125

Phe Val Ser Leu Leu Ser Ala Cys Val Tyr Gly Val Leu Tyr Ser 130 135 140

Gly Ser Phe Trp Ile His Met Leu Ser Gly Ala Ile Leu Gly Leu Ala 145 150 155 160

Trp Met Gln Ile Ala Tyr Leu Gly His Asp Ala Gly His Tyr Gln Met 165 170 175

Met Ala Thr Arg Gly Trp Asn Lys Phe Ala Gly Ile Phe Ile Gly Asn 180 185 190

Cys Ile Thr Gly Ile Ser Ile Ala Trp Trp Lys Trp Thr His Asn Ala 195 200 205

His His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His 210 215 220

Leu Pro Met Leu Ala Val Ser Ser Lys Leu Phe Asn Ser Ile Thr Ser 225 230 235 240

Val Phe Tyr Gly Arg Gln Leu Thr Phe Asp Pro Leu Ala Arg Phe Phe 245 250 255

Val Ser Tyr Gln His Tyr Leu Tyr Tyr Pro Ile Met Cys Val Ala Arg 260 265 270 Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Ile Ser Lys Arg Lys 280 275 Ile Pro Asp Arg Gly Leu Asn Ile Leu Gly Thr Leu Ile Phe Trp Thr 295 Trp Phe Pro Leu Leu Val Ser Arg Leu Pro Asn Trp Pro Glu Arg Val Ala Phe Val Leu Val Ser Phe Cys Val Thr Gly Ile Gln His Ile Gln 330 Phe Thr Leu Asn His Phe Ser Gly Asp Val Tyr Val Gly Pro Pro Lys Gly Asp Asn Trp Phe Glu Lys Gln Thr Arg Gly Thr Ile Asp Ile Ala Cys Ser Ser Trp Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu 375 Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu Arg Ser Ile Ser Pro Ile Cys Arg Glu Leu Cys Lys Lys Tyr Asn Leu Pro Tyr Val Ser Leu Ser Phe Tyr Asp Ala Asn Val Thr Thr Leu Lys Thr Leu Arg 425 Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Gln Asn 435 Leu Ala Trp Glu Ala Phe Asn Thr His Gly 455 <210> 14 <211> 35 <212> DNA <213> Artificial Sequence <223> Definition of Artificial Sequence: PCR primer for 5' of pk0011.d5 tttgcggccg caaatcaatg gaagaagcaa agaag 35 <210> 15 <211> 33 <212> DNA <213> Artificial Sequence <223> Definition of Sequence: antisense PCR primer for 3' of pk0011.d5 <400> 15 33 tttgcggccg ccaggattca cccgaaagtg ttc <210> 16 <211> 823 <212> DNA



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Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly
Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala
                         55
Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe
Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp
                                     90
Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg
                                105
            100
Val Gly
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